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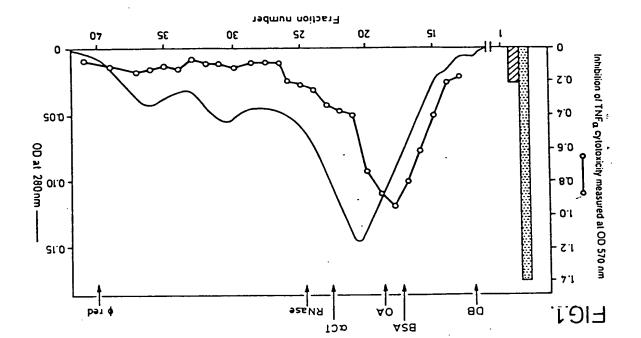
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(54) TNF-alpha inhibitors

(57). A protein having a selective tumour necrosis factor (TNF)-α inhibitory activity, but which does not block other proteins, particularly IL-1, is prepared by extraction from urine of febule patients.

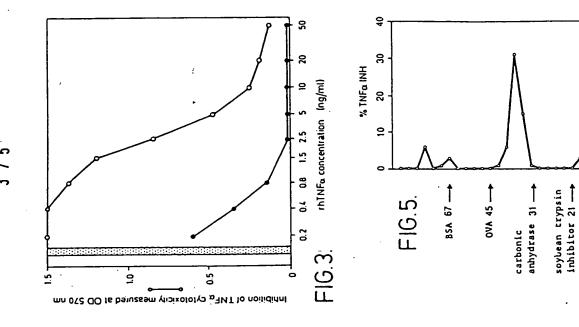
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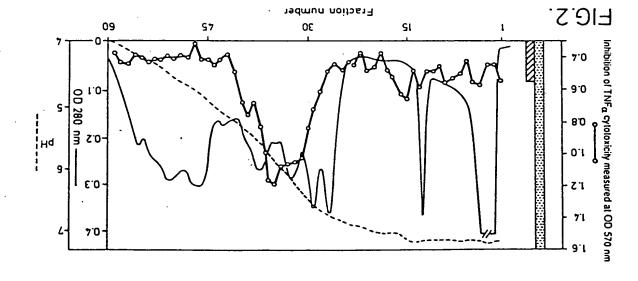




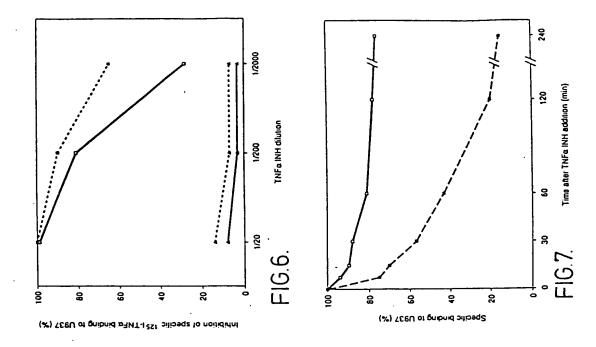
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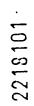


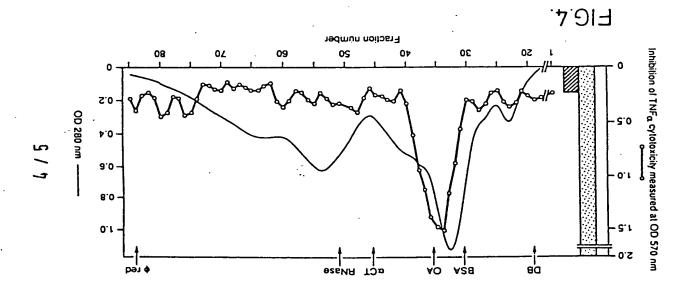
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### BIOLOGICALLY ACTIVE PROTEINS

sources, to its preparation by DNA manipulation and to the use of such protein in the treatment of conditions associated with excessive or inhibitory effect against Tumour Necrosis Factor a-mediated activity, to the isolation and purification of such a protein from natural The present invention relates to a novel protein having an unregulated INFa production.

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which accompany invasive stimuli such as bacteria, viruses, tumours and other toxins. INFB, commonly termed "lymphotoxin", is mainly produced by lymphoid cells. INFB has many activities similar to those of INFa, 림 factor a (INFa), also termed "cachectin", is mainly produced by cells 'An endotoxin-induced gerum factor that causes necrosis of tumours", but it appears to be lesa potent although this may be as a result of of the monocyte/macrophage lineage in response to "stress" signals cumour cells and inhibit their growth in culture [E. Carswell et. Tumour necrosis factor (TNF) is an activity embodied by a Natl. Acad. Sci. USA, 72, p3666 (1975)]. Tumour necrosts family of at least two proteins,  $\alpha$  and  $\beta$  , which are cytotoxic for difficulties in preparing pure  $\text{INF}\beta$  . 20

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p234 (1987)]. Elevated levela of INFa induced by, for example, tumour activities [8. Beutler et. al., "Identity of tumour necrosis factor INFa mediates and participates in a wide range of biological (1985)] sharing several of them with interleukin 1 (IL-1) [J. Le et. multiple overlapping biological activities", <u>Laboratory Invest</u>., 56, and the macrophage-secreted factor cachectin", <u>Nature</u>, 316, p552 al., "lumour necrosis factor and interleukin 1 : cytokines with

etimulation of vascular endothelial cells which release, for example, implicated as a principal mediator of endotoxic shock (septic shock) cells may lead to weight loss and cachexia and INFa has also been tissue factor) and tissue destruction (induced by, for example, can be fatal. Other biological effects of INFa include prostoglandin  $E_2$  (PGE $_2$ ) synthesis), cosquiopathy (induced by hypotension, fever (induced by stimulation of hypothalamic

production by dermal fibroblasts and synovial cells) [C. Dinarello atimulation of a series of proteinases, including collagenase

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el., "Iumour necrosis factor (cachectin) is an endogenous pyrogen al., "Cachectin/tumour necrosis factor stimulate and induces production of interleukin 1", <u>J. Exp. Med.</u>, 163, p1433 fibroblasts and synovial cells", J. Exp. Med., 162, p2163 (1985)]. collagenase and prostaglandin  $\mathsf{E}_2$  production by human dermal (1986); J. Dayer et.

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deleterious effects described above. It has been shown that passive endotoxin-Induced death, medlated by INFa antibodies [8. Beutler et. there exists, therefore, a need to develop a cachectin/INF3 Inhibitor which prevents endotoxic shock, cachexia and the other immunisation of animals against cachectin can prevent el., Nature, 316, supre]. 20

inhibitory effect against INFα-mediated activities without aignificant hereinafter identified as Tumour Necrosis factor a Inhibitor (INFa concomitant inhibition of IL-1-mediated activity. The protein is We have now identified a novel protein which has a potent INI)

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Thus in one aspect of the invention, we provide a protein which selectively inhibits tumour necrosis factor a-mediated activity.

used herein, aslective inhibition as shown by the inhibitor of in common with INF certain but not all of the biological activities of activity while lacking the ability to block other proteins which have the invention is identified as the ability to block INF-mediated INF, such as IL-1.

Invention is in a substantially homogeneous form, being substantially referably, the tumour necrosis factor  $\alpha$  inhibitor of the free from major contaminents and/or aubatantially free from other proteinaceous material. The tumour necrosis factor a inhibitor according to the invention has been found to have one or more of the following characteristics:

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(a) a molecular weight in the range 40 to 60 kDa, determined by molecular aleve chromatography;

(b) an iso-electric point (pl) in the range 5.5 to 6.1, determined by chromatofocussing;

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described by G. Nedwin et. al. "Effects of interleukin 2, interferon-γ and mitogens on the production of tumour necrosis factors a and B", <u>J.</u> although inhibition of INFa in this assay is more efficient than that competitive. The inhibitor is also an inhibitor of INF $\beta$  activity, cytotoxicity for murine 1929 cells treated with actinomycin D, as Immunol., 135, p2492 (1985). This inhibition can be overcome by (c) inhibition of the standard INF assay of differential further addition of INFa, indicating that the inhibition is of INFB;

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(d) inhibition of INF-induced  $\mathrm{PGE}_2$  release from human fibroblasts and synovial cells;

- (e) the inhibitor interferes with the binding of INFx to U937 cells (a monocytic tumour line) as evidenced by inhibition of binding of radiolabelled INFa ( $^{12}$  I-INFa);
- (f) the dissociation of a pre-formed INFa : U937 cell complex is promoted by the inhibitor in a temperature dependent manner;
- (g) the inhibitor does not degrade INF by proteolytic cleavage;
- (h) the inhibitor does not inhibit IL-1 receptor-binding activity e.g. the binding of radiolabelled IL-1 ( $^{12}$  I-IL-1a) to the murine thymoma aubline El4-6.1.

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We have found that the protein of the invention when further purified has a molecular weight of about 33000 daltons as determined by sodium dodecyl aulphate polyacrylamide gel electrophoresis (SDS PAGE).

There is thus provided as a further or alternative aspect, a protein which selectively inhibits TNFa-mediated activity which has one or more of the following characteristics:

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- (a) a molecular weight of about 33 kDa, determined by SDS PAGE;
- 20 (b) an iso-electric point (pl) in the range 5.5 to 6.1, determined by chromatofocussing;
- (c) inhibition of the standard INF assay of differential cytotoxicity for murine L929 cells treated with actinomycin D, as described by G. Nedwin et. al. "Effects of interleukin 2, interferon-γ and mitogens on the production of tumour necrosis factors a and β", 3. Immunol., 135, p2492 (1985). This inhibition can be overcome by further addition of INFα, indicating that the inhibition is

competitive. The inhibitor is also an inhibitor of INF  $\epsilon$  activity, although inhibition of INF $\alpha$  in this assay is more efficient than that of INF $\epsilon$ ;

- (d) inhibition of INF-induced PGE release from human
- fibroblasts and aynovial cells;
- (e) the inhibitor interferes with the binding of INFa to U937 cells (a monocytic tumour line) as evidenced by inhibition of binding of radiolabelled INFa ( $^{1\,\Sigma}$  I-INFa);
- (f) the dissociation of a pre-formed INFa : U937 cell complex is promoted by the inhibitor in a temperature dependent manner;

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- . (g) the inhibitor does not degrade INF by proteolytic cleavage;
- (h) the inhibitor does not inhibit IL-1 receptor-binding activity e.g. the binding of radiolabelled IL-1 ( $^{12}$  I-IL-1a) to the murine thymoma subline EL4-6.1.

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Preferably the INFa INH of the present invention has both of the characteristics (a) and (b) and one or more of the characteristics (c) to (h).

In particular, the INFa INH of the present invention has all of the characteristics (a) to (h).

The protein of the invention has an amino terminal amino acid

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sequence as follows:

Asp-Ser-Val-Cys-Pro-Gln-Cly-Lys-Tyr-Ile-HisPro-Gln-Cys-Asn-Ser-Ile

It is further believed that the next three umino acids provide a glycosylation site and that the sequence thus continues

Asn-Ser-Thr-Lys.

It will be appreciated that a INFa inhibitor according to the invention will comprise an amino acid sequence substantially corresponding to the sequence of native INFa INH and containing an amino terminal sequence substantially identical to that described

above. The sequence of a TNFa inhibitor according to the invention will thus be identical to the sequence of native INFa INH or contain one or more deletions, substitutions, insertions, inversions or additions of allelic origin or otherwise, the resulting sequence will have at least 80% and preferably 90%, homology with the sequence of native INFa INH and retain essentially the same biological properties of the protein.

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The INFa inhibitor of the invention has been demonstrated to be proteinaceous in that it is inactivated by heating in a time and temperature dependent manner is is destroyed by treatment with trypsin or promose.

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The INFa INH of the invention has also been shown to be a glycoprotein since treatment with the enzyme Endoglycosidase F reduces the molecular weight by 7 to 8 kDa.

In a further or alternative aspect of the invention there is thus provided a INFa inhibitor as defined herein, but which is in a substantially unglycosylated state.

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The inhibitors of the invention are of interest in the treatment of conditions in which it is desirable to inhibit INFa activity, for example, those which arise from the effects of INFa such as weight loss, shock, cachexia and chronic local inflammation, rheumatoid arthritis, disseminated intravascular coagulation and nephritis.

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There is thus provided as a further aspect of the invention elves inhibitor as herein defined or a pharmaceutically acceptable

INTs inhibitor as herein defined or a pharmaceutically acceptable derivative thereof for use as an active therapeutic agent, in particular, in the treatment of conditions associated with excessive or unregulated INFs production.

In a further or alternative aspect of the invention there is provided a method for the treatment of conditions associated with excessive or unregulated INFa production in a mammal including man comprising administration of an effective amount of a INFa inhibitor as herein defined or a pharmaceutically acceptable derivative thereof.

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There is also provided in a further or alternative aspect of the invention use of a INFa inhibitor as herein defined or a pharmaceutically acceptable derivative thereof for the manufacture of a medicament for the treatment of conditions associated with excessive or unregulated INFa production.

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It will be appreciated by those akilled in the art that reference herein to treatment extends to prophylaxis as well as the treatment of established conditions or symptoms.

It will be further appreciated that the amount of TNF a inhibitor of the invention required for use in treatment will vary not only with the route of administration but also with the nature of the condition being treated and the age and condition of the patient and will be ultimately at the discretion of the attendant physician or veterinarian. In general however, a suitable dose will be in the range of from about 5.0 to 500 µg per kilogram of bodyweight per day,

۰ ھ for example, in the range 30 to 300µg′kg/day, preferably in the range 50 to 150µg′kg′day.

The desired dose may conveniently be presented in a single dose or as divided doses administered at appropriate intervals, for example, as two, three, four or more sub-doses per day.

While it may be possible that, for use in therapy, a INFa inhibitor of the invention may be administered as the raw protein it is preferable to present the active protein as a pharmaceutical formulation.

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The invention further provides a pharmaceutical formulation comprising a INFa inhibitor as herein defined or a pharmaceutically acceptable derivative thereof together with one or more pharmaceutically acceptable carriers thereof and, optionally, other therapeutic and/or prophylactic ingredients. The carrier(s) must be 'acceptable' in the sense of being compatible with the ingredients of the formulation and not deleterious to the recipient thereof.

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The inhibitors according to the invention may therefore be formulated for parenteral administration (e.g. by injection, for example bolus injection or continuous infusion) and may be presented in unit dose form in ampoules, pre-filled syringes, small volume infusions or in multi-dose containers with an added preservative. The compositions may take such forms as suspensions or solutions in aqueous vehicles, and may contain formulatory agents such as suspending, stabilising and/or dispersing agents. Alternatively, the active ingredient may be in powder form obtained by aseptic isolation of sterile solid or by lyophilisation from solution, for constitution

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with a suitable vehicle, e.g. sterile, pyrogen-free water, bofore use.

The INTs inhibitor of the invention may also be used in combination with other therapeutic agents, for example, other cytokines or inhibitors thereof.

The invention thus provides, in a further aspect, a combination comprising a INFa inhibitor as herein defined or or pharmaceutically acceptable derivative thereof together with another therapeutically active agent, for example, other cytokines or inhibitors thereof.

In the proteins of the invention may be prepared by purification from natural sources and, where appropriate followed by chemical modification, or they may be prepared by conventional methods known in the art for the preparation of proteins, for example, by recombinant DNA techniques.

provided a process for producing the tumour necrosis factor a inhibitor of the invention by purification from natural sources, particularly the unine of human febrile patients. Such purification, for example, comprises the steps of concentrating the crude unine of febrile human patients, precipitating crude INFa INH from the unine and fractionating the INFa INH from the other proteins of this precipitate by one or more of, for example, fon exchange column chromatography, immunoabsorption and affinity chromatography on chromatography, immunoabsorption and affinity chromatography on immunoabsorption immunoabsorption and affinity chromatography on immunoabsorption and affinity chromatography or immunoabsorption immunoabso

The tumour necrosis factor a inhibitor of the invention is also obtainable from macrophage containing human tissue, for example,

lung lavages and extracts of human liver, from which it may be obtained by standard purification techniques such as those described above.

Natural and recombinant INFa INH produced according to the procedures described herein may be purified through a series of steps as listed above. After each of the purification steps, the presence and purity of the INFa INH may be messured in an assay of cytotoxicity in the presence of actinomycin D (acti D) using a INF-ausceptible cell line L929, as described by G. Nedwin et al., J. Immunol., 135, 10c.

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In a preferred embodiment of the process the INFa INH is initially isolated from untreated urine collected from febrile human patients (>38.5°C) devoid of urinary infections using a standard concentration technique, for example, ultrafiltration. A crude fraction may then be precipitated from the crude urine using ammonium sulphate e.g. by addition of ammonium sulphate to a concentration of 80% (w/v) at 4°C with stirring. Preferably the ammonium sulphate may be added in a stepwise manner and material precipitated at lower concentrations e.g. at 40% (w/v) discarded. The ammonium sulphate may be removed by dialysis and the resulting fraction purified to separate the INFa INH from other proteins by a variety of chromatographic methods.

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Thus, the INFa INH concentrate may be purified by ion-exchange chromatography which asparates proteins according to their differences in electrical charge, which is a reflection of the acid'base properties of the proteins. Sultable materials for anion-exchange chromatography include aminoethylcellulose derivatives, for example,

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quaternary-aminoethylcellulose (GAE-cellulose) or diethylaminoethyl- .
cellulose (CEAE-cellulose) which are widely commercially available.
The anion-exchange column should be equilibrated prior to epolying the concentrate using a suitable buffer, such as Iria-HCI, optionally containing a chelating agent such as EDIA. Bound material may be eluted from the column using a salt solution (for example, 0.8H sodium chloride made up with the equilibration buffer).

The pooled active fractions from the anion-exchange chromatograph chromatography. Sultable materials for cation-exchange chromatograph 10 include derivatives of cellulose such as carboxymethyl (CM) cellulose or Sulphopropyl Sepharose (Pharmacia, Uppsala, Sweden). The column should be equilibrated with a suitable buffer, such as sodium acetate and bound material may be eluted with the equilibration buffer containing, for example, 0.5M sodium chloride.

chromatography on bound recombinant human INFa (rhINFa), coupled to a suitable matrix, for example, Mini-Leak Agarose (Kem En Jec, Biotechnology Corp., Dermark). The column should be buffered using, for instance, a phosphate buffer (e.g. 0.8H potassium phosphate pH for instance, a phosphate buffer (e.g. 0.8H potassium phosphate pH ethanolamine-HCl pH 8.5 buffer. The column should be equilibrated with a suitable buffer, for example, Iria-HCl optionally containing sodium chloride and the INFa INH eluted with an acidic (pH 3.5) glycine buffer. The eluted fractions should immediately be balanced to pH 7.0 by the addition of, for example, Iria base.

The active pooled fractions are preferably lyophilised prior to the final purification step of reverse-phase FPLC (fast protein liquid

cells are yeart cells, <u>E. coll</u> cells and animal cells.

Expression of a protein having tunour necrosis fector inhibitor activity is achieved by culture of the transformed host. cells in a suitable growth medium. Normally such a medium will contain a source of nitrogen such as ammonium sulphate, a source of carbon and energy such as glucose or glycerol, trace elements and factors essential to growth of the particular host cells. The precise culture conditions will be dependent upon the chosen host; thus, for example, in the case of E. coli submerged aerobic fermentation is preferred, preferably at about 37°C.

In addition, expression may be induced, for example, by the addition of an inducer or the use of inducing conditions for the promoter system being used in the expression vector.

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Depending upon the host, the INFa inhibitor may be produced as granular inclusion bodies which can be recovered, after cell lysis, by differential centrifugation; these can be solubilised by conventional methods and purified by the methods described herein for purification of urinary INFa INH. Alternatively, the INFa inhibitor may be in solution in the cytosol, secreted into the periplasmic space or conveniently secreted into the culture medium.

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The host cells are transformed by recombinant DNA molecules which comprise a DNA sequence encoding for a INFa inhibitor which has been inserted into an expression vector.

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Such expression vectors may consist of segments of chromosomal, non-chromosomal and synthetic DNA sequences, such as various known derivatives of SV-40 and known bacterial plasmics, for example, "natural" plasmids such as ColEl, pSCIOI or pRSF2124 and

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phare DKAs, or "artificiai" plasmius (constructed in vitro) such as a pBR322, pi89 or p41153. Phage DVAs include, for example, the numerous derivatives of phage lambda and other DNA phages, for example H13, and other filamentous single-stranded DNA phages. Vectors useful in yeasts include the 2p plasmid, and those useful in eukaryotic cells such as animal cells include those containing SV-40 adenovirus and retrovirus.

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Such expression vectors may also be characterised by at least one expression control sequence which may be operatively linked to the Inhibitor DNA sequence such that it controls and regulates the expression of the cloned DNA sequence. Examples of useful expression control sequences include the lac, trp, tac and trc systems, major operator and promoter regions of phage A (such as the P<sub>L</sub> promoter under the control of the thermolabile ts cl857 repressor), the control region of fd cost protein, the glycolytic promoters of yeast (e.g. the promoter for 3-phosphoglycerate kinase), the promoters of yeast acid phosphatase (e.g. Pho 5), the promoters of yeast a-mating factors, and promoters derived from polyoma, adenovirus, retrovirus, and simian virus.

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In addition, such expression vectors may possess various sites for insertion of the INFa inhibitor DNA sequences of this invention. These sites are characterised by the specific restriction endonuclease which cleaves them. Such cleavage sites are well recognised by those skilled in the art. The expression vector, and in particular the site chosen therein for insertion of a selected DNA fragment and its operative linking to an expression control sequence, is determined by a variety of factors including the number of sites susceptible to a

given restriction enzyme, the size of the protein to be expressed, contemination or binding of the protein to be expressed by host ceil proteins which may be difficult to remove during purification, the location of start/stop codons, and other factors recognised by those skilled in the art. Thus the choice of a vector and insertion site for a DNA sequence is determined by a balance of these factors, not all selections being equally effective for a given case.

Likewise, not all host/vector combinations will function with equal efficiency in expressing the DNA sequences of this invention.

The selection is made, depending upon a variety of factors including compatability of the host and vector, ease of recovery of the desired protein, expression characteristics of the DNA sequences and the expression control sequences operatively linked to them, or eny necessary post-expression modifications of the desired protein.

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The DNA sequences of the invention which on expression code for proteins with INFa inhibitor activity may be isolated by screening various DNA libraries for such DNA sequences using a series of DNA probes. The DNA probes may be prepared from the purified natural protein which is used as a source of amino acid sequence data. The purified natural protein may be prepared, for example, from febrile human unine as described above. Degenerate DNA sequences coding for various portions and fragments of the amino acid sequence, e.g. in combination with Lathe probes, are used to design the DNA probes.

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Thus, various DNA libraries are screened for DNA sequences coding for the INFa inhibitors of the invention. Such libraries include chromosomal gene banks and cDNA or DNA libraries prepared from cell lines or tissue that are demonstrated to produce INFa

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inhibitors, such as alveolar macrophages or liver tissue. Screening may be direct immune expression, for example in Agill or sittler systems, or, in the case where a TMI INH producing cell is identified, by identification of TMF INH specific mNA by direct expression in Xenopus occytes.

A variety of conventional cloning and selection techniques may be used to locate and identify DNA sequences 'ALL' incode on expression in an appropriate eukaryotic or prokaryotic host for the INFa inhibitors of this invention. These selected DNA sequences may themselves be used as probes to select other DNA sequences coding for INF1 inhibitors or may be used in appropriate recombinant DNA molecules to transform appropriate eukaryotic or prokearyotic hosts for the production of INF1 INH encoded by them.

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The invention includes within its scope single and double stranded DNA sequences encoding for INFa INIB, vectors containing such sequences suitable for transformation of a host organism and host cells transformed with such DNA sequences.

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According to a further aspect of the present invention we provide a protein with selective INFa inhibitor activity produced by expression of a host transformed with a DNA sequence encoding for such a INFa inhibitor protein. INF inhibitors of the invention which are prepared by the expression of a DNA sequence encoding such inhibitors in a transformed host will thus be identical to the sequence of native INFa INH or contain one or more deletions, substitutions, insertions, inversions or additions of allelic origin or otherwise, the resulting sequence will have at least 80% and preferably 90% homology with the sequence of native INFa INH and retain essentially the same biological

include en N-terminal methionine. Also, for example, the DNA senuence encoding DNA sequence or aid secretion, maturation or purification of ģ of the lavention coding for INFa INH may be fused in an expression intra-or extra-cellularly by known techniques or the INFa INH may vector to a portion of a DNA sequence coding for a sukariotic or the INFa INH from the host; the fused polypeptide may be removed prokaryotic polypeptide to assist the expression of the INFa INI properties. In perticuler, a INF inhibitor of the invertion may together with the fused polypeptide.

INHs can then be employed, after purification, in the pharmaceutical prokaryotic hosts transformed with DNA sequences encoding for INFa the INFa INNs produced by culturing of the eukaryotic and compositions of this invention. It will be appreciated that, when produced by animal cells, the unglycosylated state. In addition, the glycosylated protein may be substantially deglycosylated by techniques known in the art, for INFa INH of the invention will be a glycoprotein. Prokaryotic expression systems will, however, produce the protein in an example, by the use of endoglycosidase enzymes.

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All temperatures are in  $^{
m 0}$ C and all percentage concentrations are  $^{
m w'v}.$ the following non-limiting Examples illustrate the invention. 20

### INFo Inhibition Assay

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D) corresponded to 100% inhibition, whereas the OD from cells cultured the percentage of INFa INH activity in the fractions described (0D) values from murine L929 cells stimuloted by actinomycin D (acti in the Examples was determined by assuming that the optical density

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Blochem., 152, p515 (1985). Thus the percentage of INFa inhibition in human INFa (rhINFa) produced in <u>E. coli</u> as described by A. Harmenout with actinomycin D and  ${\rm INFa}$  corresponded to a maximal cell mostality el., "Molecular cloning and expression of human tumour necrosis of O% INFa inhibition. The INFa used in the assay was recombinent factor and comparison with mouse tumour necrosis factor", Eur. J. the assay of cytotoxicity was calculated accord.∵; to formula (I)

Percentage of INFa INH activity =

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### Exemple 1

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Purification of Uninary INFa INH

## Concentration of Protein from Human Urine

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Buffering from small-cell carcinoma, one from malignant histocytosis, Human uning (15 litres) was freshly obtained from a pool of one from polymyocitis and one from sepsis. All were highly febrile five patients prior to any treatment. Iwo of the patients were concentrated at 40 on an Amicon ultrafiltration hollow fibre (>38.5 $^{\circ}$ C) and devoid of uninary infections. The unine was apparatus, with a molecular size cut-off of about 5 kDa

## b) Precipitation of Protein from Human Urine

ammonium sulphate. A pellet was obtained by centrifugation which was sodium chlorids. The ammonium sulphate was removed by dialysis at 4º The concentrated unine pool was saturated with solid ensorice aulphate by adding the sulphate slowly with constant stirring at 4° Ë resuspended in 150 ml of 20 mM sodium phosphate (pM 7.2) and 150 using 10 m/ Iris-HCl pH 7.4, 2 m/ EDIA and 5 m/ benzamidine HCl. aupernatant adjusted to 80% saturation with addition of further until an ammonium sulphate concentration of 40% was reached. precipitate was removed by centrifugation, discarded, and the 9

### c) Identification of INFa INH Activity

fraction, total inhibition of the cytotoxic effect induced by rhiNfa presence of actinomycln D. At a 1:20 dilution of the semi-purified measured in the presence of actinomycin D alone  $(00_{57\mu nm} = 1.5)$ . the semi-purified fraction of Example 1(b) was tested in cytoxicity assay with the INF-susceptible cell-line L929 in the was observed so that the OD<sub>570nm</sub> value was identical to that

control value of rhINFa at a final concentration of 0.2 ng/ml measured furthermore, inhibitory activity was observed in dilutions of that 50% of inhibition was observed at a dilution of approximately the fraction of up to 1:160 on cells  $(00_{\rm 570}{\rm nm}$  = 0.83) whereas the in the presence of actinomycin D was lower  $(00_{57\,\text{Unm}}$  = 0.73), so The INFa .INH had no effect on cell viability when tested without actinomycin D. 1:100 (0D<sub>570nm</sub> = 1.10).

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Comparison of the Effect of TVFs 15% on INF a and 6 Induced Cutotoxicity

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are shown in Table 1, below, and demonstrate that the inhibitor of the invention does have some inhibitory effect on INF; mediated cytoxicity semi-purified fraction from Example 1(b) was tested at 1:20, 1:50 and ranging from approximately 50% down to 2% of the INFa inhibition with INFa or INFB cytokine, and in the absence of inhibitor. The results A cytotoxicity assay was conducted using the INF-susceptible 1:80 dilutions. Control tests were performed in the absence of the concentration of INFa or INFB to induce the cytotoxic effect. The cell-line L929 in the presence of actinomycin D using a range of increasing INFA concentration. 2

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TABLE 1

Final Concentration of TMF (a or 3) (pg/ml) Added to Actionycin-D	form of Cytokine added to	Final S-290 L929 c	Dilution of Sephany Inhibitory Fraction cells (OD <sub>57</sub> unm <sup>2</sup> )	tion of Sephaceyl Sitory Fraction o (ODs.yunm.	50 oc.
	CSIIS	None	1,20	1,′50	1,80
0	0	>1.90	11.90	>1.90	>1.90
10	: :	Ç	NÇ.	S	Q
	<b>6</b> 0.	1.16	1.66	1.46	1.39
. 20	8	1.30	>1.90	1.72	1.68
	<b>ന</b> .	1.02	1.51	1.21	1.22
50	ಕ	1.02	>1.90	1.72	1.68
	ന.	0.65	1.03	0.84	0.68
100	8	0.73	1.78	1.69	1.51
	ør.	0.37	0.71	0.59	0.52
250	ø	0.38	1.70	1.70	1.33
	ar	0.24	0.44	0.31	0.24
\$00	ช	0.30	1.52	1.49	1.11
	<b>62</b> .	0.17	0.30	0.26	0.18
1 250		0.19	1.09	1.10	0.76
	œ.	0.11	0.13	0.19	0.13
2 500	ø	90.0	1.03	0.80	0.47
	<b>6</b> 2.	S	ON	S.	QN

#### Example 2

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### Gel Filtration of Uninery INFr INH

fraction (20 mg, 0.8 ml) was applied to the caluma and sluted with the cm) (Pharmacia, Uppsala, Sweden) equilibrated in 50 mM Iris-HCl buffer eluted from the gel in a single peak. The inhibitory activity showed filtration chromatography at 4 $^{\rm 0}$  on a Sephacryl S-200 column (0.9 x 60 (pH7.4) containing 100 mH sodium chlorids. A sample of the protein collected and tested for INFa INH activity. The INFa INH activity same buffer at a flow-rate of 5.4 ml/hr. fractions (1.35 ml) were The semi-purified TMFa INH of Example 1(b) was purified by gel an apparent molecular weight of 40 to 60 kDa (aee Figure 1). 유

#### Example 3

### Chromatofocussing of Urinary INFa INH

chromatofocussed at 4 $^{0}$  on a Hono-P pre-packed column (HR 5/20, 5  $\times$  200 mm) (Pharmacia, Uppsala, Sweden) equilibrated in 25 mM Bis-Iris buffer 74/iminodiacetic acid at pH 4.0. Column fractions (1 ml) were tested actual pH of each column fraction was determined with a pH meter, the bulk of the INFa INH activity being contained in the eluted fractions Switzerland). A sample of the protein fraction of Example 1(b) (30 cytotoxicity assay in the presence of actinomycin D (l $\mu g/m l$ ). The at 1:10 dilution for their effect in the rhINF $\alpha$  (0.2 ng/ml) mg) was applied to the column and eluted with a polybuffer adjusted to pH 7.1 with imidodiacetic acid (Fluka, Buchs, The semi-purified INFs INH of Example 1(b) was 20 25 15

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ä between pH 5.5 and 6.1 (see Figure 2). This is equivalent to the of the INFa INH protein.

#### Example 4

## Ion-Exchange Chromatography of Urinary INFa INH

buffer pH 8.0, containing 2mil EDIA. Bound material was eluted from the fractions (8.0ml) were collected, tested for INFa INH activity and the column with the equilibration byffer containing 0.8% sodium chloride. anion-exchange chromatography at  $4^{\circ}$  on a DEAE Sephadex column (2.6  $\times$ Inhibitory fractions were pooled (160ml) and dialysed against 10mi 20cm) (Pharmacia, Uppsala, Sweden) equilibrated in 10m/ Tris-HCl The semi-purified INFa INH of Example 1(b) was purified by sodium acetate buffer pH 5.0 (4 x 2 litres).

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Inhibitory fractions were pooled and concentrated 20-fold on an Amicon chromatography at  $4^0$  on a Sulphopropyl-Sephadex column (0.8 x 15 cm) ultrafiltration apparatus with a molecular size cut-off of about 10 buffer pH 5.0. Bound material was eluted from the column with the (Pharmacia, Uppsala, Sweden) equilibrated in 10mM sodium acetate equilibration buffer containing 0.5H sodium chloride. Fractions (7.5ml) were collected, tested for INFa INH activity and the the INFa INH was further purified by catton-exchange kDa.

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### Example 5

### Gel Filtration of Unionry INF: INH 25

filtration chromatography at  $4^{\circ}$  on a Sephacryl S-200 column (2.6 imes 100The INFa INH concentrate from Example 4 was purified by gel

A sample of the protein with the equilibration buffer at a flow-rate of 27ml/hour. frections raction from Example 4 (200mg) was applied to the column and eluted cm) (Phermecia, Uppsala, Sweden) equilionated with Somi Tris-101 ph The column was calibrated with Jaxtran blue (DB), 2000 kDa; bovine serum albumin (BSA), 67 kDa; ovalbumin (OA), 43 kDa; a -chymotrypsinogen-A (aCI), 25 kDa; and (9.0ml) were collected, tested for INFa INH activity and ribonuclease A (RNase), 13.5 kDa, as shown in Figure 4. 7.4 Euffer containing 100mil sodium chloride. inhibitory fractions were pooled.

#### Example 6

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## Affinity Chromatography of Urinary INFa INH

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collected, immediately adjusted to pH 7.0 by addition of IM.Iris (5 to A Tafa effinity column was prepared by coupling recombinant human containing 100m! sodium chloride (3 x 50ml). A sample of the INFa INH fractions from Example 5 (15ml) was applied to the column and eluted ouffer pH 8.5. The gel was washed with 50m/1 Iris-HCl pH 7.4 buffer INFa (1.0mg) to Mini Leak Agarose (Kem En Tec, Biotechnology Corp., Denmark) in 0.811 potassium phosphate buffer pH 8.6. The remaining active groups were blocked by incubation in 0.1H ethanolamine-HCl with a G.2H glycine-HCl pH 3.5 buffer. Fractions (1.0ml) were Op1) and tested for INFa INH activity.

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#### Example 7

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# Reverse Phase FPLC Chromatography of Urinary INFa INH

The INFa INH fractions from Example 6 were lyophilised, dissolved in 0.1% trifiluoroacetic acid (2.0ml) and loaded onto a ProRPC

with a 0 to 100% acetonitrile gradient in 0.1% trifluoroacetic acid at equilibrated in 0.1% trifluoroacetic acid. Bound material was eluted a flow rate of 0.3ml/minute. To each fraction (0.75ml) 0.5M ammonium bicarbonate (1 $\mathfrak Q_1$ 1) was added and the eluted material was lyophilised. reverse-phase FPLC column (5 x 20 cm) (Pharmacia, Uppsala, Sweden)

containing this activity were diasolved in 10mM Tris-HC1 pH 7.4 buffer described by U. Laemmli et al., Nature, 277, p 680 (1970). The INFa INH was found to elute with a molecular weight of 33 kDa (see Figure mg/ml of rhINFa. The activity directed against rhINFa migrated with an apparent molecular weight identical to the 33 kDa band on the gel Samples run under non-reducing conditions were tested for INFa INH activity at 1:10 dilution on L929 cells in the presence of 0.15 he reverse-phase FPLC chromatography revealed one major peak polyacrylamide gel electrophoresis (505 PAGE) using the method corresponding to INFa INH activity. The lyophilised fractions containing 2mM EDIA and analysed by sodium dodecyl aulphate run under reducing conditions.

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### Example 8

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## Protein Sequencing of Uninary INFa INH

N,N-diisopropylethylamine acetate and acetonitrile prior to injection. Applied Blosystems Model 477A protein sequencer. Fractions from the conditioned sequencer filter. The protein was analysed with an aequencing cycles were evaporated to dryness and resuspended in The INFa INH frection isolated from the reverse-phase FPLC chromatography was concentrated in vacuo and spotted onto into an HPLC column for residue identification.

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- 27 -

Tyr-lle-His-Pro-Gin-Cys-Asn-Ser-Ile. It is further believed that the identified and have the sequence : Asp-Ser-Val-Cys-Pro-Gin-Gly-Lyssignificantly homogeneous to any protein sequence contained in the next three amino acids provide a glycosylation aite and that the sequence thus continues Asn-Ser-Thr-Lys. This sequence is not The first 17 amino acid residues of the N-terminal were

#### Example 9

NBRF Protein Sequence database (November 1988).

- Demonstration that INFa INH is a Protein 2
- a) Time and Temperature Dependency

750 The Sephacryl S-200 purified  $INF\alpha$  INH of Example 2, obtained minutes and, by comparison with untreated samples, the percentage of by bulking the tubes of the active fractions, was heated at  $56^{
m o}$ , and 95°. The INFa INH activity was measured after 10, 20 and 60 INFa INH activity was calculated according to formula (1).

decreases in a time- and temperature-dependent manner.

results shown in Table 2 below demonstrate that the INFa INH activity

•		Heat inactivation	
	Temperature ( <sup>0</sup> C)	Itme (min)	Percentage of INFa INH activity
•	56	10 20 60	100 100 93
	. 52	10 20 60	60 26 15
10	\$6	. 10 20 60	27 10 13

### b) Susceptibility to Irypsin Digestion

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Irypsin (500μg) (51gma, 5t. Louis, NO) in 0.2M Iris-HCl buffer (pH 8.0) containing lam calcium chloride was added to the pooled fractions of Sephacryl 5-200 purified urinary INFα INH of Example 2 and incubated at 37°C for 4 hours. Another measure of trypsin (500μg) was added and digestion continued for a further 20 hours, at which time the reaction was terminated by addition of soybean trypsin inhibitor (2mg) (5igma, 5t. Louis, NO). The percentage of INFα INH inhibitory activity of the trypsin digest and the control was determined at a 1:20 final dilution of the pool of bulked fractions on L929 cells stimulated by rhINFα in the presence of actinomycin D, according to formula (1). The results are shown in Table 3 below.

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Tr	Trypsin inactivation	
Conditions	Pscentage of INfa INH activity	OC 570nm
Buffer alone	0	0.71
Irypsin + soybean trypsin inhibitor in buffer	· ·	0.70
Partially purified Sephadex 5-200 urine	61	1.46
Partially purified Sephadex:5-200 urine digested by trypsin	23	1.03

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### c) Treatment with Urea

The Sephacryl S-200 purified INFa, INH of Example 2 was adjusted to 2H urea and extensively dialysed at 40 against phosphate buffered saline (PBS) containing 2H urea. Dialysis against PBS was repeated prior to the bloassay. INFa INH activity was found to be unaffected which indicates that inhibitory activity is not mediated by a molecule of low molecular weight bound to a protein.

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### Example 10

## Demonstration of Competitive Inhibition

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The Sephacryl S-200 purified INFa INH of Example 2 was tested at a 1:10 dilution against increasing amounts of rhINFa on L929 cells. An inverse correlation between the amount of rhINFa present in the assay and the degree of inhibition was observed (see Figure 3). Thus, the inhibitor, activity is competitively overcome by increasing concentrations of rhINFa.

### Example 11

# Inhibition of INFa-Mediated PCE, Production by Dermal Fibroblasts

Human dermal fibroblasts were seeded at a concentration of  $2.0~\mathrm{x}$ cells/well and cultured for 48 hours. Cells were then stimulated hours of incubation, PGE $_2$  production was measured in the supernatants ᆡ with DMEM buffer supplemented with 10% FCS as a control. Cells were 5mg/ml, and the effect of the INFa INH from Example 5 was studied at three dilutions (1:20, 1:50 and 1:80) in the above buffer. After 72 also atimulated with rhINR at concentrations ranging from 0.5 to by radioimmunoassay using a PGE $_2$  antiserum [see J M Dayer et al., Clin. Invest., 67, pl386 (1979)].

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The results are shown in Table 4, below, and demonstrate that the At ability of rhTNFa to atimulate PGE $_2$  production by dermal fibroblasts мвь inhibited by the addition of INFa INH at all three dilutions. 1:80 dilution of INFa INH the inhibitory activity was partially overcome by increasing rhINFa concentrations.

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Concentration of rhIMF PGC2 production by human dermal fibroblasts (ng/ml)	PG£ <sub>2</sub> production	by human dermal	fibroblasts (ng/m	=
on human fibroblasts	Dilutio	Dilution of TRF INH on fibroblasts	fibroblasts	
(1ш/6d)	none	1:80	1:50	1:20
0	50.6 ± 7.4	88.8 ± 5.6	103.0 ± 8.9	111.0 ± 9.4
005	160.0 ± 14.1	126.0 2 9.3	115.9 ± 6.61	113.9 ± 7,1
2,000	331.7 ± 28.4	217.2 ± 10.7	156.7 ± 10.7	115.3 ± 21.3
2,000	381.7 ± 19.6	257.2 ± 13.7	253.1 ± 21.2	221.6 ± 16.0

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concentrations of rhillfu. PGE2 production by cultured human dermal fibroblasts was measured after ihree different experiments were carried out with the same strzin of fitroblasts. Duffer or INFo 1484 was incubated at various dilutions in the presence or absence of various three days. Values represent triplicate means of the three cultures  $\pm$  S(H (u,j)

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### Example 12

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### rhINFa Binding Inhibition Assay

was cultured at  $4^0$  for 2 hours in a culture medium ( $200\mu\mathrm{J}$ ) comprising The apecific activity of [1251]-INFa was 2.2xl04 cpm/ng and produced Fraker and Speck Jr., Blochem. Blophys. Res. Comm., 80, p849 (1978). (100µg/ml), penicillin (100U/ml), 1.0% glutamine and 10% foetal calf The human macrophage cell line U937 in aliquots of  $10^6\,$  cells RPMI:1640 (Gibco, Paialey, Scotland) aupplemented with streptomycin Recombinant human INFa was lodinated by using the lodogen method of single band with a molecular weight of 17 kDa when analysed by SDS  $[^{12}$  ]]-INFa. Binding inhibition was performed by the addition of serum, and additionally containing 0.04% sodium azide and 0.5ng various dilutions of INF INH (1:20, 1:200 and 1:2000).

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from the bound [12 I]-INFa by centrifugation through an oil mixture as Non-specific binding was measured in the presence of a 100-fold excess of unlabelled rhINF $\alpha$ , and free readloactivity was separated described by Robb et al., J. Exp. Med., 154, p1455 (1981). Cell Sweden), and the percentage of binding inhibition was determined bound [ $^{1.5}$  I]—INR was measured in a gamma counter (LKB, Bromma, according to formula (II)

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Percentage of binding inhibition =

### Example 13

# Effect of TN5. INH on $[^{1.5}$ $_{ m I}]$ -TNFa Bindina to U937 Cells

u937 cells were preincubated for I hour at  $20^{6}$  in the culture medium of Example 12, in the presence of either [ $^{12}$  I]-INFa alone or [ $^{12}$ I]-INFa with a 100-fold excess of unlabelled rhINFa. The U937 cells were then washed with phosphate buffered saline (3 x 50ml) at 4 $^{9}$ . The cells incubated in [ $^{125}$ I]-INFa alone were divided into four batches and incubated with INFa INH from Example 5 (1:20, 1:200 and 1:2000 dilutions) and with buffer alone, respectively.

The specific binding of [<sup>1,2</sup>]-INFa to U937 cells was found to be inhibited at 40 by 100%, 80% and 35% by the three dilutions of INFa INH, 1:20, 1:200 and 1:2000, respectively (see Figure 5). The control batch which lacked INFa INH showed no inhibitory activity. The binding inhibition of the two weaker dilutions was found to be increased to 90% and 60% when [<sup>1,2</sup>5]-INFa was preincubated with INFa INH at dilutions 1:200 and 1:2000, respectively, prior to cell addition.

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The experiment was repeated using the cells preincubated in the presence of [ $^{12}$  I]-TNF $_{\alpha}$  and a 100-fold excess of unlabelled TNF $_{\alpha}$  so that the percentage of binding inhibition could be corrected for nonspecific binding.

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### Example 14

## Dissociation of a Pre-formed INFa:U937 Complex

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1937 cells were preincubated for 1 hour in the presence of [18 I]-INfa as described in Example 12. The cells were washed and incubated at either  $4^0$  or 370 in the presence or absence of INfa INII

from Example 5. Cell-surface bound [1.25]j-iNfa was found to dissociate faster in the presence of INfa INH than in its absence and this was found to occur in a time- and temperature-dependent manner (see Figure 6).

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### Example 15

# Demonstration that INFa INH is not Protecuytic for the INFa

[12 I]-INFa was incubated at  $20^{0}$  for 1 hour in the presence of 3 different dilutions of INF INH (1:20, 1:200 and 1:2000) and in the presence of buffer alone. When analysed by SOS PAGE and autoradiography the [125I]-INFa was found to migrate as a single band both in the absence and presence of INFa INHI showing the inhibitor to have no proteolytic effect.

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### Examole 16

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# Effect of INFc INH on IL-1 Receptor Binding Activity

The activity of INFa INH from Example 5 was tested in the IL-1/LAF (lymphocyte activating factor) assay when induced by IL-la or IL-lg [this assay is described by P. Seckinger et al., J. Immunol., 139, p1541 (1987) for an IL-l inhibitor protein]. A dose response of [3H]-thymidine incorporation (corresponding to thymocyte proliferation) was observed in up to 200pg/ml concentrations of both IL-la and IL-lg. Addition of INFa INH at levels observed to inhibit rhINFa did not have any significant effect on the IL-l-induced thymocyte proliferation, proving inhibition to be specific for INFa only.

The results obtained wre illustrated by reference to the accompanying drawings, in which:-

Figure 1 - shows the uninary TNFa INH activity profile of Sephacryl S-200 gel filtration. Column fractions (Iml) were tested at 1:10 dilution for effect in the rhINFa (1.0ng/ml) cytotoxicity assay in the presence of actinomycin D (1.0µg/ml) (o——o). The line (——) represents 00280nm of the fractions. Bars represent cell lysis measured by dye uptake in response to actinomycin D ([3]) and to actinomycin D plus hrINFa ([3]) without unine. The molecular weight markers are dextran blue (DB), bovine serum albumin (BSA), ovalbumin (OA), a-chymotrypsinogen (aCI), ribonuclease A (RNase) and phenol red (o-red).

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chromatofocussing on a Mono-P column. Column fractions (lml) were tested at 1:10 dilution for effect in the rhINFa (0.2ng/ml) cytotoxicity assay in the presence of actinomycin D (1.0 µg/ml) (0...). The line (....) represents OD<sub>280nm</sub> of the fractions, and (.....) represents their pH. The bars are as described for Figure 1.

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circles (o——o) represent OD<sub>570nm</sub> measured in the presence of actinomycin D, rhINFa with INFa INH; closed circles (•——•) represent OQ<sub>70nm</sub> measured in the presence of actinomycin and rhINFa only and the bar (国) represents OD<sub>570nm</sub> in the presence of actinomycin 0 (1.0µg/ml) alone.

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Figure 4 - shows the elution profile of Sepacryl S-200 gel , filtration with purified INFa INH from Example 5. Column fractions , (9ml) were sterilized and tested at 1:50 dilution against rhINFa , (1.0mg/ml) in the presence of actinomycin D (1.0µg/ml) in the presence of actinomycin D (1.0µg/ml) in the L929 cytotoxicity assay (o----o). The line (-----) represents 00<sub>280</sub>nm of the fractions. The bars are as defined for Figure 1.

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Example 7. SDS PAGE was performed as described by U. Laemmil et al., Nature, 277, loc cit. Samples were loaded onto 15% polyacrylamide gel with a 3% stacking gel and gels were sliver-stained as described by C. Herril et al., Proc. Natl. Acad. Sci. USA, 76, p4335 (1979). Samples run under non-reducing conditions were tested for biological activity by cutting 2mm slices from the gel and eluting the proteins by overnight incubation in 10mM Tris-HCl pH 7.4 containing 2mM EDIA (total volume 200µ1). Fractions were tested at 1:10 dilution on L929 cells in the presence of rhINFa (0.15ng/ml).

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complex. U937 cells were pre-incubated with [\$^{12}\$I]=Infa, washed and incubated with INfa INH as described in Example 14, at either 40 (\$^{-1}\$I] or 370 (\$^{-1}\$I]. At the time indicated, cell associated radioactivity was measured and percentage specific binding determined. On the graph, the value obtained from the control without the inhibitor has been substracted from the values obtained at the two temperatures, thus, 100% corresponds to the value obtained without the addition of INFa INH.

CLAIMS

1. A protein which inhibits tumour necrosis factor (INF) a-mediated activity but does not block other proteins which have in common with INF certain but not all of the biological activities of INF.

A protein which selectively inhibits tumour necrosis factor a-mediated activity and having one or more of the following characteristics:

(a) a molecular weight in the range 40 to 60 kDa, determined by molecular sieve chromatography;

(b) an iso-electric point (pl) in the range 5.5 to 6.1, determined by chromatofocussing; (c) inhibition of the standard INF assay of differential cytotoxicity for murine L929 cells treated with actinomycin D;

(d) inhibition of TNF-induced PGE $_{\rm 2}$  release from human fibroblasts and synovial cells;

(e) the inhibitor interferes with the binding of INFa to U937 cells (a monocytic tumour line) as evidenced by inhibition of binding of radiolabelled INFa ( $^{125}$  I-INFa);

(f) the dissociation of a pre-formed INFa : U937 cell complex is promoted by the inhibitor in a temperature dependent manner;

(g) the inhibitor does not degrade INF by proteolytic cleavage;

(h) the inhibitor does not inhibit IL-1 receptor-binding activity e.g. the binding of radiolobelled IL-1 ( $^{12}$ )-IL-1 $_u$ ) to the murine thymoma subline EL4-6.1.

A protein which selectively inhibits tumour necrosis factor a-mediated activity and having one or more of the following

(a) a molecular weight of about 33 kDa, determined by SDS

characteristics:

PAGE;

(b) an iso-electric point (pI) in the range 5.5 to 6.1, determined by chromatofocussing; (c) inhibition of the standard INF assay of differential cytotoxicity for murine L929 cells treated with actinomycin D;

(d) inhibition of TNF-induced PGE $_2$  release from human

fibroblasts and synovial cells;

(e) the inhibitor interferes with the binding of INFa to U937 cells (a monocytic tumour line) as evidenced by inhibition of binding of radiolabelled INFa ( $^{125}$  I-INFa);

(f) the diasoclation of a pre-formed INFa : U937 cell complex is promoted by the inhibitor in a temperature dependent manner;

(g) the inhibitor does not degrade INF by proteolytic

cleavage;

(h) the inhibitor does not inhibit IL-1 receptor-binding activity e.g. the binding of radiolabelled IL-1 ( $^{125}$ I-IL-1a) to the murine thymoma aubline [L4-6.1.

4. A protein as claimed in either of claims 2 or 3 having the properties
 properties (a) and (b) together with one or more of the properties
 (c) to (h).

5. A protein as claimed in either of claims 2 or 3 having all of the properties (a) to (h).

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6. A protein according to any one of claims 1 to 5 which corresponds to a naturally occurring  $TNF\alpha$  inhibitor.

7. A protein according to any one of claims 1 to 6 having an amino

terminal amino acid sequence as follows :

Asp-Ser-Val-Cys-Pro-Cln-Lys-Tyr-lle-His-Pro-Gln-Cys-Asn-Ser-lle.

Gly missing, Yell. A

> 8. A protein according to claim 7 having an amino terminal amino acid sequence as follows :

Asp-Ser-Val-Cya-Pro-Cln-Cly-Lya-Tyr-Ile-His-Pro-Cln-Cys-Asn-Ser-Ile-Asn-Ser-Thr-Lys, 9. A protein according to any one of claims 1 to 8 and in which the amino acid sequence contains one or more deletions, substitutions, insertions, inversions or additions of allelic origin or otherwise, the resulting sequence having at least 80% homology with the parent protein and retaining essentially the same hiological properties as the parent protein.

10. A protein according to claim 9 having at least 90% homology with the parent protein.

- A protein according to any one of claims 1 to 10 in a substantially homogeneous form.
- 12. A protein according to any one of claims 1 to 11 which is recombinant protein.
- A protein according to any one of claims 1 to 12 which is a glycosylated protein.
- 14. A protein according to any one of claims 1 to 12 which is in a substantially unglycosylated state.
- 15. An exogeneous DNA comprising a nucleotide sequence coding for a protein as defined in any one of claims 1 to 11.
- 16. A dONA comprising a nucleotide sequence coding for a protein according to any one of claims 1 to 11.
- A recombinant expression vector comprising DNA according to either of claims 15 or 16.
- 18. A host cell transformed with an expression vector according to claim 17.

- 19. A method of producing a INFa INH protein which comprises culturing a cell according to claim 18 and isolating the INFa INH protein.
- A recombinant protein produced according to the method of claim
   19.
- 21. A method for the preparation of a INFa INH protein comprising the steps of :
- (a) concentration of unine from febrile patients;
- (b) ammonium sulphate precipitation;
- (c) anion—exchange chomatography;
- (d) cation-exchange chromatography;
- (e) gel filtration;
- (f) affinity chrometography; and
- (g) reverse phase FPLC.
- 22. A protein characterised in that it is substantially identical to the protein obtained by the method of claim 21.
- 23. A pharmaceutical formulation comprising a INFa inhibitor as defined in any one of claims 1 to 14 or claim 22 or a pharmaceutically acceptable derivative thereof and a pharmaceutically acceptable carrier therefor.

24. A protein as defined in any one of claims 1 to 14 or claim 22 for use in therapy.

25. A pharmaceutical formulation for use in the manufacture of a medicament for the treatment of conditions associated with excessive or unregulated INFa production, wherein said formulation comprises a INFa inhibitor as defined in any one of claims 1 to 14 or claim 22 or or a pharmaceutically acceptable derivative thereof.

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